

SEQUENCE LISTING

<110> XIANG, JINHUA
WÜNSCHMANN, SABINA
SCHMIDT, WARREN
STAPLETON, JACK T.

<120> FULL-LENGTH GB VIRUS C (HEPATITIS G VIRUS) RNA TRANSCRIPTS ARE INFECTIOUS
IN PRIMARY CD4 POSITIVE T CELLS

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<141> 2000-11-27

<150> 60/195,597
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<170> PatentIn Ver. 2.1

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Ser Leu Arg Arg Leu Ala Arg Val Val Glu Cys Cys Val Met Ala Gly
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Gly Gly Arg Val Thr Ala Ala Arg Phe Thr Arg Pro Trp Thr Gln Val
1125 1130 1135

Pro Thr Asp Ala Lys Thr Thr Glu Pro Pro Pro Val Pro Ala Lys
1140 1145 1150

Gly Val Phe Lys Glu Ala Pro Leu Phe Met Pro Thr Gly Ala Gly Lys
1155 1160 1165

Ser Thr Arg Val Pro Leu Glu Tyr Gly Asn Met Gly His Lys Val Leu
1170 1175 1180

Ile Leu Asn Pro Ser Val Ala Thr Val Arg Ala Met Gly Pro Tyr Met
1185 1190 1195 1200

Glu Arg Leu Ala Gly Lys His Pro Ser Ile Tyr Cys Gly His Asp Thr
1205 1210 1215

Thr Ala Phe Thr Arg Ile Thr Asp Ser Pro Leu Thr Tyr Ser Thr Tyr
1220 1225 1230

Gly Arg Phe Leu Ala Asn Pro Arg Gln Met Leu Arg Gly Val Ser Val
1235 1240 1245

Val Ile Cys Asp Glu Cys His Ser His Asp Ser Thr Val Leu Leu Gly
1250 1255 1260

Ile Gly Arg Val Arg Glu Leu Ala Arg Gly Cys Gly Val Gln Leu Val
1265 1270 1275 1280

Leu Tyr Ala Thr Ala Thr Pro Pro Gly Ser Pro Met Thr Gln His Pro
1285 1290 1295

Ser Ile Ile Glu Thr Lys Leu Asp Val Gly Glu Ile Pro Phe Tyr Gly
1300 1305 1310

His Gly Ile Pro Leu Glu Arg Met Arg Thr Gly Arg His Leu Val Phe
1315 1320 1325

Cys His Ser Lys Ala Glu Cys Glu Arg Leu Ala Gly Gln Phe Ser Ala
1330 1335 1340

Arg Gly Val Asn Ala Ile Ala Tyr Tyr Arg Gly Lys Asp Ser Ser Ile
1345 1350 1355 1360

Ile Lys Asp Gly Asp Leu Val Val Cys Ala Thr Asp Ala Leu Ser Thr
1365 1370 1375

Gly Tyr Thr Gly Asn Phe Asp Ser Val Thr Asp Cys Gly Leu Val Val
1380 1385 1390

Glu Glu Val Val Glu Val Thr Leu Asp Pro Thr Ile Thr Ile Ser Leu
 1395 1400 1405
 Arg Thr Val Pro Ala Ser Ala Glu Leu Ser Met Gln Arg Arg Gly Arg
 1410 1415 1420
 Thr Gly Arg Gly Arg Ser Gly Arg Tyr Tyr Ala Gly Val Gly Lys
 1425 1430 1435 1440
 Ala Pro Ala Gly Val Val Arg Ser Gly Pro Val Trp Ser Ala Val Glu
 1445 1450 1455
 Ala Gly Val Thr Trp Tyr Gly Met Glu Pro Asp Leu Thr Ala Asn Leu
 1460 1465 1470
 Leu Arg Leu Tyr Asp Asn Cys Pro Tyr Thr Ala Ala Val Ala Ala Asp
 1475 1480 1485
 Ile Gly Glu Ala Ala Val Phe Phe Ser Gly Leu Ala Pro Leu Arg Met
 1490 1495 1500
 His Pro Asp Val Ser Trp Ala Lys Val Arg Gly Val Asn Trp Pro Phe
 1505 1510 1515 1520
 Leu Val Gly Val Gln Arg Thr Met Cys Arg Glu Thr Leu Ser Pro Gly
 1525 1530 1535
 Pro Ser Asp Asp Pro Gln Trp Ala Gly Leu Lys Gly Pro Asn Pro Val
 1540 1545 1550
 Pro Leu Leu Leu Arg Trp Gly Asn Asp Leu Pro Ser Lys Val Ala Gly
 1555 1560 1565
 His His Ile Val Asp Asp Leu Val Arg Arg Leu Gly Val Ala Glu Gly
 1570 1575 1580
 Tyr Val Arg Cys Asp Ala Gly Pro Ile Leu Met Val Gly Leu Ala Ile
 1585 1590 1595 1600
 Ala Gly Gly Met Ile Tyr Ala Ser Tyr Thr Gly Ser Leu Val Val Val
 1605 1610 1615
 Thr Asp Trp Asp Val Lys Gly Gly Ser Pro Leu Tyr Arg His Gly
 1620 1625 1630
 Asp Gln Ala Thr Pro Gln Pro Val Val Gln Val Pro Pro Val Asp His
 1635 1640 1645
 Arg Pro Gly Gly Glu Ser Ala Pro Ser Asp Ala Asn Thr Val Thr Asp
 1650 1655 1660
 Ala Val Ala Ala Ile Gln Val Asp Cys Asp Trp Ser Val Met Thr Leu
 1665 1670 1675 1680
 Ser Ile Gly Glu Val Leu Ser Leu Ala Gln Ala Lys Thr Ala Glu Ala
 1685 1690 1695

Tyr Ala Ala Thr Thr Lys Trp Leu Ala Gly Cys Tyr Thr Gly Thr Arg
 1700 1705 1710
 Ala Val Pro Thr Val Ser Ile Val Asp Lys Leu Phe Ala Gly Gly Trp
 1715 1720 1725
 Ala Ala Val Val Gly His Cys His Ser Val Ile Ala Ala Ala Val Ala
 1730 1735 1740
 Ala Tyr Gly Ala Ser Arg Ser Pro Pro Leu Ala Ala Ala Ala Ser Tyr
 1745 1750 1755 1760
 Leu Met Gly Leu Gly Val Gly Gly Asn Ala Gln Thr Arg Leu Ala Ser
 1765 1770 1775
 Ala Leu Leu Leu Gly Ala Ala Gly Thr Ala Leu Gly Thr Pro Val Val
 1780 1785 1790
 Gly Leu Thr Met Ala Gly Ala Phe Met Gly Ser Ala Ser Val Ser Pro
 1795 1800 1805
 Ser Leu Val Thr Ile Leu Leu Gly Ala Val Gly Gly Trp Glu Gly Val
 1810 1815 1820
 Val Asn Ala Ala Ser Leu Val Phe Asp Phe Met Ala Gly Lys Leu Ser
 1825 1830 1835 1840
 Ser Glu Asp Leu Trp Tyr Ala Ile Pro Val Leu Thr Ser Pro Gly Ala
 1845 1850 1855
 Gly Leu Ala Gly Ile Ala Leu Gly Leu Val Leu Tyr Ser Ala Asn Asn
 1860 1865 1870
 Ser Gly Thr Thr Trp Leu Asn Arg Leu Leu Thr Thr Leu Pro Arg
 1875 1880 1885
 Ser Ser Cys Ile Pro Asp Ser Tyr Phe Gln Gln Ala Asp Tyr Cys Asp
 1890 1895 1900
 Lys Val Ser Ala Val Leu Arg Arg Leu Ser Leu Thr Arg Thr Val Val
 1905 1910 1915 1920
 Ala Leu Val Asn Arg Glu Pro Lys Val Asp Glu Val Gln Val Gly Tyr
 1925 1930 1935
 Val Trp Asp Leu Trp Glu Trp Ile Met Arg Gln Val Arg Met Val Met
 1940 1945 1950
 Ala Arg Leu Arg Ala Leu Cys Pro Val Val Ser Leu Pro Leu Trp His
 1955 1960 1965
 Cys Gly Glu Gly Trp Ser Gly Glu Trp Leu Leu Asp Gly His Val Glu
 1970 1975 1980
 Ser Arg Cys Leu Cys Gly Cys Val Ile Thr Gly Asp Val Leu Asn Gly
 1985 1990 1995 2000

Gln Leu Lys Asp Pro Val Tyr Ser Thr Lys Leu Cys Arg His Tyr Trp
 2005 2010 2015
 Met Gly Thr Val Pro Val Asn Met Leu Gly Tyr Gly Glu Thr Ser Pro
 2020 2025 2030
 Leu Leu Ala Ser Asp Thr Pro Lys Val Val Pro Phe Gly Thr Ser Gly
 2035 2040 2045
 Trp Ala Glu Val Val Val Thr Pro Thr His Val Val Ile Arg Arg Thr
 2050 2055 2060
 Ser Ala Tyr Lys Leu Leu Arg Gln Gln Ile Leu Ser Ala Ala Val Ala
 2065 2070 2075 2080
 Glu Pro Tyr Tyr Val Asp Gly Ile Pro Val Ser Trp Asp Ala Asp Ala
 2085 2090 2095
 Arg Ala Pro Ala Met Val Tyr Gly Pro Gly Gln Ser Val Thr Ile Asp
 2100 2105 2110
 Gly Glu Arg Tyr Thr Leu Pro His Gln Leu Arg Leu Arg Asn Val Ala
 2115 2120 2125
 Pro Ser Glu Val Ser Ser Glu Val Ser Ile Asp Ile Gly Thr Glu Thr
 2130 2135 2140
 Glu Asp Ser Glu Leu Thr Glu Ala Asp Leu Pro Pro Ala Ala Ala
 2145 2150 2155 2160
 Leu Gln Ala Ile Glu Asn Ala Ala Arg Ile Leu Glu Pro His Ile Asp
 2165 2170 2175
 Val Ile Met Glu Asp Cys Ser Thr Pro Ser Leu Cys Gly Ser Ser Arg
 2180 2185 2190
 Glu Met Pro Val Trp Gly Glu Asp Ile Pro Arg Thr Pro Ser Pro Ala
 2195 2200 2205
 Leu Ile Ser Val Thr Glu Ser Ser Pro Asp Glu Lys Thr Pro Ser Val
 2210 2215 2220
 Ser Ser Ser Gln Glu Asp Thr Pro Ser Ser Asp Ser Phe Glu Val Ile
 2225 2230 2235 2240
 Gln Glu Ser Glu Thr Ala Glu Gly Glu Glu Ser Val Phe Asn Val Ala
 2245 2250 2255
 Leu Ser Val Leu Lys Ala Leu Phe Pro Gln Ser Asp Ala Thr Arg Lys
 2260 2265 2270
 Leu Thr Val Lys Met Ser Cys Cys Val Glu Lys Ser Val Thr Arg Phe
 2275 2280 2285
 Phe Ser Leu Gly Leu Thr Val Ala Asp Val Ala Ser Leu Cys Glu Met
 2290 2295 2300

Glu Ile Gln Asn His Thr Ala Tyr Cys Asp Lys Val Arg Thr Pro Leu
2305 2310 2315 2320

Glu Leu Gln Val Gly Cys Leu Val Gly Asn Glu Leu Thr Phe Glu Cys
2325 2330 2335

Asp Lys Cys Glu Ala Arg Gln Glu Thr Leu Ala Ser Phe Ser Tyr Ile
2340 2345 2350

Trp Ser Gly Val Pro Leu Thr Arg Ala Thr Pro Ala Lys Pro Pro Val
2355 2360 2365

Val Arg Pro Val Gly Ser Leu Leu Val Ala Asp Thr Thr Lys Val Tyr
2370 2375 2380

Val Thr Asn Pro Asp Asn Val Gly Arg Arg Val Asp Lys Val Thr Phe
2385 2390 2395 2400

Trp Arg Ala Pro Arg Val His Asp Lys Phe Leu Val Asp Ser Ile Glu
2405 2410 2415

Arg Ala Lys Arg Ala Ala Gln Ala Cys Leu Ser Met Gly Tyr Thr Tyr
2420 2425 2430

Glu Glu Ala Ile Arg Thr Val Arg Pro His Ala Ala Met Gly Trp Gly
2435 2440 2445

Ser Lys Val Ser Val Lys Asp Leu Ala Thr Pro Ala Gly Lys Met Ala
2450 2455 2460

Val His Asp Arg Leu Gln Glu Ile Leu Glu Gly Thr Pro Val Pro Phe
2465 2470 2475 2480

Thr Leu Thr Val Lys Lys Glu Val Phe Phe Lys Asp Arg Lys Glu Glu
2485 2490 2495

Lys Ala Pro Arg Leu Ile Val Phe Pro Pro Leu Asp Phe Arg Ile Ala
2500 2505 2510

Glu Lys Leu Ile Leu Gly Asp Pro Gly Arg Val Ala Lys Ala Val Leu
2515 2520 2525

Gly Gly Ala Tyr Ala Phe Gln Tyr Thr Pro Asn Gln Arg Ile Arg Glu
2530 2535 2540

Met Leu Lys Leu Trp Glu Ser Lys Lys Thr Pro Cys Ala Ile Cys Val
2545 2550 2555 2560

Asp Ala Thr Cys Phe Asp Ser Ser Ile Thr Glu Glu Asp Val Ala Leu
2565 2570 2575

Glu Thr Glu Leu Tyr Ala Leu Ala Ser Asp His Pro Glu Trp Val Arg
2580 2585 2590

Ala Leu Gly Lys Tyr Tyr Ala Ser Gly Thr Met Val Thr Pro Glu Gly
2595 2600 2605

Val Pro Val Gly Glu Arg Tyr Cys Arg Ser Ser Gly Val Leu Thr Thr
2610 2615 2620

Ser Ala Ser Asn Cys Leu Thr Cys Tyr Ile Lys Val Lys Ala Ala Cys
2625 2630 2635 2640

Glu Arg Val Gly Leu Lys Asn Val Ser Leu Leu Ile Ala Gly Asp Asp
2645 2650 2655

Cys Leu Ile Ile Cys Glu Arg Pro Val Cys Asp Pro Ser Asp Ala Leu
2660 2665 2670

Gly Arg Ala Leu Ala Ser Tyr Gly Tyr Ala Cys Glu Pro Ser Tyr His
2675 2680 2685

Ala Ser Leu Asp Thr Ala Pro Phe Cys Ser Thr Trp Leu Ala Glu Cys
2690 2695 2700

Asn Ala Asp Gly Lys Arg His Phe Phe Leu Thr Thr Asp Phe Arg Arg
2705 2710 2715 2720

Pro Leu Ala Arg Met Ser Ser Glu Tyr Ser Asp Pro Met Ala Ser Ala
2725 2730 2735

Ile Gly Tyr Ile Leu Leu Tyr Pro Trp His Pro Ile Thr Arg Trp Val
2740 2745 2750

Ile Ile Pro His Val Leu Thr Cys Ala Phe Arg Gly Gly Thr Pro
2755 2760 2765

Ser Asp Pro Val Trp Cys Gln Val His Gly Asn Tyr Tyr Lys Phe Pro
2770 2775 2780

Leu Asp Lys Leu Pro Asn Ile Ile Val Ala Leu His Gly Pro Ala Ala
2785 2790 2795 2800

Leu Arg Val Thr Ala Asp Thr Thr Lys Thr Lys Met Glu Ala Gly Lys
2805 2810 2815

Val Leu Ser Asp Leu Lys Leu Pro Gly Leu Ala Val His Arg Lys Lys
2820 2825 2830

Ala Gly Ala Leu Arg Thr Arg Met Leu Arg Ser Arg Gly Trp Ala Glu
2835 2840 2845

Leu Ala Arg Gly Leu Leu Trp Arg Pro Gly Leu Arg Leu Pro Pro Pro
2850 2855 2860

Glu Ile Ala Gly Ile Pro Gly Gly Phe Pro Leu Ser Pro Pro Tyr Met
2865 2870 2875 2880

Gly Val Val His Gln Leu Asp Phe Thr Ser Gln Arg Ser Arg Trp Arg
2885 2890 2895

Trp Leu Gly Phe Leu Ala Leu Leu Ile Val Ala Leu Phe Gly
2900 2905 2910

